

Timothy Triche

List of Publications by Year in descending order

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Version: 2024-02-01

18
papers

5,822
citations

840776

11
h-index

940533

16
g-index

20
all docs

20
docs citations

20
times ranked

10739
citing authors

#	ARTICLE	IF	CITATIONS
1	Non-Invasive Profiling of Advanced Prostate Cancer via Multi-Parametric Liquid Biopsy and Radiomic Analysis. <i>International Journal of Molecular Sciences</i> , 2022, 23, 2571.	4.1	8
2	Abstract 295: Superior efficacy of targeted nanoparticle therapy for brain tumors. <i>Cancer Research</i> , 2022, 82, 295-295.	0.9	0
3	Abstract 1636: A new invasive zebrafish model of Ewing sarcoma reveals EWSR1-FLI1-driven dysregulation of heparan sulfate proteoglycan metabolism in developing tumors. <i>Cancer Research</i> , 2022, 82, 1636-1636.	0.9	0
4	Baseline circulating tumor cell (CTC) count as a prognostic marker of PSA response and progression in metastatic castrate sensitive prostate cancer (mCSPC): Results from SWOG S1216, a phase III randomized trial of androgen deprivation plus orteronel (cyp17 inhibitor) or bicalutamide.. <i>Journal of Clinical Oncology</i> , 2020, 38, 5506-5506.	1.6	4
5	A phase 1 study of azacitidine combined with chemotherapy in childhood leukemia: a report from the TACL consortium. <i>Blood</i> , 2018, 131, 1145-1148.	1.4	44
6	The molecular landscape of pediatric acute myeloid leukemia reveals recurrent structural alterations and age-specific mutational interactions. <i>Nature Medicine</i> , 2018, 24, 103-112.	30.7	525
7	Tumor Variant Identification That Accounts for the Unique Molecular Landscape of Pediatric Malignancies. <i>JNCI Cancer Spectrum</i> , 2018, 2, pky079.	2.9	8
8	Response: Making "perfect" the enemy of good. <i>Blood</i> , 2017, 130, 1168-1169.	1.4	2
9	Molecular Profiling Reveals Biologically Discrete Subsets and Pathways of Progression in Diffuse Glioma. <i>Cell</i> , 2016, 164, 550-563.	28.9	1,695
10	Comprehensive, Integrative Genomic Analysis of Diffuse Lower-Grade Gliomas. <i>New England Journal of Medicine</i> , 2015, 372, 2481-2498.	27.0	2,582
11	The Genomic Landscape of the Ewing Sarcoma Family of Tumors Reveals Recurrent STAG2 Mutation. <i>PLoS Genetics</i> , 2014, 10, e1004475.	3.5	335
12	Genome-Wide Characterization of Circulating Tumor Cells Identifies Novel Prognostic Genomic Alterations in Systemic Melanoma Metastasis. <i>Clinical Chemistry</i> , 2014, 60, 873-885.	3.2	47
13	Canine tumor cross-species genomics uncovers targets linked to osteosarcoma progression. <i>BMC Genomics</i> , 2009, 10, 625.	2.8	228
14	Selective and Titratable Effects On AML Genome Wide Methylation Patterning, Transcription and Clonogenicity Using Very Low Concentrations of 5-Aza-2'deoxyctidine.. <i>Blood</i> , 2009, 114, 2385-2385.	1.4	0
15	Expression Profiles of Osteosarcoma That Can Predict Response to Chemotherapy. <i>Cancer Research</i> , 2005, 65, 8142-8150.	0.9	132
16	Infectomics: genomics and proteomics of microbial infections. <i>Functional and Integrative Genomics</i> , 2002, 1, 331-344.	3.5	44
17	Genetic Determinants of Coxsackievirus B3 Pathogenesis. <i>Annals of the New York Academy of Sciences</i> , 2002, 975, 169-179.	3.8	19
18	Novel genomic imbalances in embryonal rhabdomyosarcoma revealed by comparative genomic hybridization and fluorescence in situ hybridization: An Intergroup Rhabdomyosarcoma Study. <i>Genes Chromosomes and Cancer</i> , 2000, 27, 337-344.	2.8	141